SEQUENCE LISTING

SEQ ID NO:1

human CNG3B amino acid sequence

MFKSLTKVNKVKPIGENNENEQSSRRNEEGSHPSNQSQQTTAQEENKGEEKSLKTKSTPVTS
EEPHTNIQDKLSKKNSSGDLTTNPDPQNAAEPTGTVPEQKEMDPGKEGPNSPQNKPPAAPVI
NEYADAQLHNLVKRMRQRTALYKKKLVEGDLSSPEASPQTAKPTAVPPVKESDDKPTEHYYR
LLWFKVKKMPLTEYLKRIKLPNSIDSYTDRLYLLWLLLVTLAYNWNCWFIPLRLVFPYQTAD
NIHYWLIADIICDIIYLYDMLFIQPRLQFVRGGDIIVDSNELRKHYRTSTKFQLDVASIIPF
DICYLFFGFNPMFRANRMLKYTSFFEFNHHLESIMDKAYIYRVIRTTGYLLFILHINACVYY
WASNYEGIGTTRWVYDGEGNEYLRCYYWAVRTLITIGGLPEPQTLFEIVFQLLNFFSGVFVF
SSLIGQMRDVIGAATANQNYFRACMDDTIAYMNNYSIPKLVQKRVRTWYEYTWDSQRMLDES
DLLKTLPTTVQLALAIDVNFSIISKVDLFKGCDTQMIYDMLLRLKSVLYLPGDFVCKKGEIG
KEMYIIKHGEVQVLGGPDGTKVLVTLKAGSVFGEISLLAAGGGNRRTANVVAHGFANLLTLD
KKTLQEILVHYPDSERILMKKARVLLKQKAKTAEATPPRKDLALLFPPKEETPKLFKTLLGG
TGKASLARLLKLKREQAAQKKENSEGGEEEGKENEDKQKENEDKQKENEDKGKENEDKDKGR
EPEEKPLDRPECTASPIAVEEEPHSVRRTVLPRGTSRQSLIISMAPSAEGGEEVLTIEVKEK
AKQ

SEQ ID NO:2

complete human CNG3B nucleotide sequence

CATTTCTCTACCTTAAGGCACAGTCATAAATACAGAGGGTTTTCAGAACCACCTCAGAGAAGATG
TTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAGGAGAGAACAATGAGAATGAACAAAG
TTCTCGTCGGAATGAAGAAGGCTCTCACCCAAGTAATCAGTCTCAGCAAACCACAGCACAGGAAG
AAAACAAAGGTGAAGAGAAATCTCTCAAAACCAAGTCAACTCCAGTCACGTCTGAAGAGCCACAC
ACCAACATACAAGACAAACTCTCCAAGAAAAATTCCTCTGGAGATCTGACCACAAACCCTGACCC
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CTACACAACCTGGTGAAAAGAATGCGTCAAAGAACAGCCCTCTACAAGAAAAAGTTGGTAGAGGG
AGATCTCTCCTCACCCGAAGCCAGCCCACAAACTGCAAAGCCCACGGCTGTACCACCAGTAAAAG
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ATCATCTACCTTTATGATATGCTATTTATCCAGCCCAGACTCCAGTTTGTAAGAGGAGGAGACAT
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CATCAATAATACCATTTGATATTTGCTACCTCTTCTTTGGGTTTAATCCAATGTTTAGAGCAAAT
AGGATGTTAAAGTACACTTCATTTTTTGAATTTAATCATCACCTAGAGTCTATAATGGACAAAGC
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CAAAGAAAAGGCTAAGCAATAAATGTTTGATTATCTTTAGATGTGATATAGCTAGTTCCCAAAGT
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GAAAGGCAAATCCCTAGCTTAGTTTCTAGGACTTATCTGAGAGTGTGATTTCATGCAGTGGTAAT
AAGAAGATTATTAAAAGCAAAAAAAAAAAAAAAAAAA

SEQ ID NO:3

human CNG3B coding sequence

ATGTTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAGGAGAGAACAATGAGAATGAACA
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CAGCTACACAACCTGGTGAAAAGAATGCGTCAAAGAACAGCCCTCTACAAGAAAAAGTTGGTAGA
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AAGAAAGCGATGATAAGCCAACAGAACATTACTACAGGCTGTTGTGGTTCAAAGTCAAAAAGATG
CCTTTAACAGAGTACTTAAAGCGAATTAAACTTCCAAACAGCATAGATTCATACACAGATCGACT
CTATCTCCTGTGGCTCTTGCTTGTCACTCTTGCCTATAACTGGAACTGCTGGTTTATACCACTGC
GCCTCGTCTTCCCATATCAAACCGCAGACAACATACACTACTGGCTTATTGCGGACATCATATGT
GATATCATCTACCTTTATGATATGCTATTTATCCAGCCCAGACTCCAGTTTGTAAGAGGAGGAGA
CATAATAGTGGATTCAAATGAGCTAAGGAAACACTACAGGACTTCTACAAAATTTCAGTTGGATG
TCGCATCAATAATACCATTTGATATTTGCTACCTCTTCTTTGGGTTTAATCCAATGTTTAGAGCA
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TGTGGTGGCCCACGGGTTTGCCAATCTTTTAACTCTAGACAAAAAGACCCTCCAAGAAATTCTAG
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AAGACCGCAGAAGCAACCCCTCCAAGAAAAGATCTTGCCCTCCTCTTCCCACCGAAAGAAGAGAC
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TGAAGATAAAGATAAAGGAAGAGAGCCAGAAGAGAAGCCACTGGACAGACCTGAATGTACAGCAA
GTCCTATTGCAGTGGAGGAAGAACCCCACTCAGTTAGAAGGACAGTTTTACCCAGAGGGACTTCT
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SEQ ID NO:4

Oligo 1 (sense strand primer)

TCTATCTCCTGTGGCTCTTGCTTGTC

SEQ ID NO:5

Oligo 2 (antisense strand primer)

GAGTCTGGGCTGGATAAATAGCATATC

SEQ ID NO:6

Oligo 3 (sense strand primer)

AGGAATTGGCACTACTAGATGGGTG

SEQ ID NO:7

Oligo 4 (antisense strand primer)

TTCATGAGGATCCTTTCAGAATCTGG

SEQ ID NO:8

Oligo 5 (sense strand primer)

GGAAACCGTCGAACTGCCAATGTGGT

SEQ ID NO:9

Oligo 6 (sense strand primer)

CGGGTTTGCCAATCTTTTAACTCTAGAC

SEQ ID NO:10

Oligo 7 (antisense strand primer)

GTCCGCAATAAGCCAGTAGTGTATG

SEQ ID NO:11

Oligo 8 (sense strand primer)

TGACAAGCTTCCGCCATGTTTAAATCGCTGACAAAAGTC

SEQ ID NO:12

Oligo 9 (antisense strand primer)

TGACGAATTCTCCCAGCATGTCGTTTCCCCTCGTTAA